

SCORE Search Results Details for Application 10552515 and Search Result 20090316_112516_us-10-552-515-7.ra1.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
Page	List	Overview	FAQ	Suggestions

This page gives you Search Results detail for the Application 10552515 and Search Result 20090316_112516_us-10-552-515-7.ra1.

[Go Back to previous page](#)

GenCore version 6.3

Copyright (c) 1993 - 2009 Bioccceleration Ltd.

OM protein - protein search, using sw model

Run on: March 17, 2009, 05:01:40 ; Search time 2 Seconds
(without alignments)
1258.128 Million cell updates/sec

Title: US-10-552-515-7
Perfect score: 40
Sequence: 1 ILILSKIYV 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1316349 seqs, 215321474 residues

Total number of hits satisfying chosen parameters: 1316349

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*
1: /ABSS/Data/CRF/ptodata/1/iaa/5_COMB.pep:*
2: /ABSS/Data/CRF/ptodata/1/iaa/6_COMB.pep:*
3: /ABSS/Data/CRF/ptodata/1/iaa/7_COMB.pep:*
4: /ABSS/Data/CRF/ptodata/1/iaa/H_COMB.pep:*
5: /ABSS/Data/CRF/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /ABSS/Data/CRF/ptodata/1/iaa/RE_COMB.pep:*
7: /ABSS/Data/CRF/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

Result No.	Score	Query Match	Length	DB	ID	Description
1	32	80.0	58	3	US-10-703-032-145414	Sequence 145414,
2	32	80.0	59	3	US-10-703-032-142763	Sequence 142763,
3	32	80.0	281	3	US-10-703-032-166070	Sequence 166070,
4	32	80.0	366	2	US-09-248-796A-17343	Sequence 17343, A
5	31	77.5	53	3	US-09-450-969-5111	Sequence 5111, Ap
6	31	77.5	53	3	US-10-724-972B-5111	Sequence 5111, Ap
7	31	77.5	167	2	US-09-270-767-35863	Sequence 35863, A
8	31	77.5	167	2	US-09-270-767-51080	Sequence 51080, A
9	31	77.5	184	2	US-09-248-796A-14687	Sequence 14687, A
10	31	77.5	289	3	US-10-369-493-7235	Sequence 7235, Ap
11	31	77.5	312	2	US-09-270-767-35094	Sequence 35094, A
12	31	77.5	312	2	US-09-270-767-50311	Sequence 50311, A
13	31	77.5	312	3	US-10-369-493-4476	Sequence 4476, Ap
14	31	77.5	758	3	US-10-149-310-260	Sequence 260, App
15	31	77.5	883	2	US-09-248-796A-20980	Sequence 20980, A
16	30	75.0	46	3	US-10-100-683-10450	Sequence 10450, A
17	30	75.0	46	3	US-11-001-793-10450	Sequence 10450, A
18	30	75.0	57	3	US-10-100-683-7848	Sequence 7848, Ap
19	30	75.0	57	3	US-11-001-793-7848	Sequence 7848, Ap
20	30	75.0	70	3	US-10-703-032-177589	Sequence 177589,
21	30	75.0	71	2	US-09-107-532A-6034	Sequence 6034, Ap
22	30	75.0	78	3	US-10-703-032-149742	Sequence 149742,
23	30	75.0	82	2	US-09-248-796A-17471	Sequence 17471, A
24	30	75.0	154	3	US-09-540-209B-8991	Sequence 8991, Ap
25	30	75.0	182	2	US-09-621-976-4098	Sequence 4098, Ap
26	30	75.0	182	3	US-10-664-025A-4098	Sequence 4098, Ap
27	30	75.0	235	3	US-09-979-932A-642	Sequence 642, App
28	30	75.0	235	3	US-09-979-932A-813	Sequence 813, App
29	30	75.0	237	2	US-09-270-767-58463	Sequence 58463, A
30	30	75.0	262	2	US-09-270-767-43128	Sequence 43128, A
31	30	75.0	319	3	US-09-540-209B-6073	Sequence 6073, Ap
32	30	75.0	327	2	US-09-543-681A-8331	Sequence 8331, Ap
33	30	75.0	394	3	US-09-876-997-259	Sequence 259, App
34	30	75.0	394	3	US-10-108-260A-4310	Sequence 4310, Ap
35	30	75.0	394	3	US-10-643-836-259	Sequence 259, App
36	30	75.0	531	2	US-09-134-001C-3574	Sequence 3574, Ap
37	30	75.0	531	3	US-09-450-969-4004	Sequence 4004, Ap
38	30	75.0	531	3	US-10-724-972B-4004	Sequence 4004, Ap
39	30	75.0	531	3	US-10-902-441B-3574	Sequence 3574, Ap
40	30	75.0	743	2	US-09-248-796A-17817	Sequence 17817, A
41	30	75.0	861	2	US-09-538-092-809	Sequence 809, App
42	30	75.0	1770	3	US-10-361-522-8	Sequence 8, Appli
43	29	72.5	41	3	US-09-450-969-5433	Sequence 5433, Ap
44	29	72.5	41	3	US-10-724-972B-5433	Sequence 5433, Ap
45	29	72.5	81	2	US-09-248-796A-22424	Sequence 22424, A

ALIGNMENTS

RESULT 1

US-10-703-032-145414

; Sequence 145414, Application US/10703032

```

; Patent No. 7214786
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Andersen, Scott E.
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Conner, Timothy W.
; APPLICANT: Cao, Yongwei
; APPLICANT: Masucci, James D.
; APPLICANT: Zhou, Yihua
; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR APPLICATION NUMBER: 10/020,338
; PRIOR FILING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 145414
; LENGTH: 58
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_TA_39832.pep
US-10-703-032-145414

```

```

Query Match          80.0%; Score 32; DB 3; Length 58;
Best Local Similarity 44.4%; Pred. No. 31;
Matches      4; Conservative      5; Mismatches      0; Indels      0; Gaps      0;

```

```

Qy      1 ILILSKIYV 9
          :|:|:|:|:
Db      47 VLVLNKLYI 55

```

RESULT 2

```

US-10-703-032-142763
; Sequence 142763, Application US/10703032
; Patent No. 7214786
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Andersen, Scott E.
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Conner, Timothy W.
; APPLICANT: Cao, Yongwei
; APPLICANT: Masucci, James D.
; APPLICANT: Zhou, Yihua
; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR APPLICATION NUMBER: 10/020,338
; PRIOR FILING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 142763

```

```

;   LENGTH: 59
;   TYPE: PRT
;   ORGANISM: Triticum aestivum
;   FEATURE:
;   OTHER INFORMATION: Clone ID: PAT_TA_37181.pep
US-10-703-032-142763

```

```

Query Match           80.0%;   Score 32;   DB 3;   Length 59;
Best Local Similarity 100.0%;   Pred. No. 31;
Matches      7;   Conservative      0;   Mismatches      0;   Indels      0;   Gaps      0;

```

```

Qy      2 LILSKIY 8
        |||
Db      35 LILSKIY 41

```

RESULT 3

```

US-10-703-032-166070
; Sequence 166070, Application US/10703032
; Patent No. 7214786
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Andersen, Scott E.
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Conner, Timothy W.
; APPLICANT: Cao, Yongwei
; APPLICANT: Masucci, James D.
; APPLICANT: Zhou, Yihua
; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR APPLICATION NUMBER: 10/020,338
; PRIOR FILING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 166070
; LENGTH: 281
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(281)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_TA_60488.pep
US-10-703-032-166070

```

```

Query Match           80.0%;   Score 32;   DB 3;   Length 281;
Best Local Similarity 75.0%;   Pred. No. 1.6e+02;
Matches      6;   Conservative      2;   Mismatches      0;   Indels      0;   Gaps      0;

```

```

Qy      1 ILILSKIY 8
        ::|||
Db      100 VVILSKIY 107

```

RESULT 4

US-09-248-796A-17343

; Sequence 17343, Application US/09248796A

; Patent No. 6747137

; GENERAL INFORMATION:

; APPLICANT: Keith Weinstock et al

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS

; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: 107196.132

; CURRENT APPLICATION NUMBER: US/09/248,796A

; CURRENT FILING DATE: 1999-02-12

; PRIOR APPLICATION NUMBER: US 60/074,725

; PRIOR FILING DATE: 1998-02-13

; PRIOR APPLICATION NUMBER: US 60/096,409

; PRIOR FILING DATE: 1998-08-13

; NUMBER OF SEQ ID NOS: 28208

; SEQ ID NO 17343

; LENGTH: 366

; TYPE: PRT

; ORGANISM: Candida albicans

US-09-248-796A-17343

Query Match 80.0%; Score 32; DB 2; Length 366;

Best Local Similarity 75.0%; Pred. No. 2.1e+02;

Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ILILSKIY 8

|:|||||:

Db 104 IVILSKVY 111

RESULT 5

US-09-450-969-5111

; Sequence 5111, Application US/09450969

; Patent No. 7060458

; GENERAL INFORMATION:

; APPLICANT: Lynn Doucette-Stamm et al.

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS

; TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: PATH99-09A

; CURRENT APPLICATION NUMBER: US/09/450,969

; CURRENT FILING DATE: 1999-11-29

; NUMBER OF SEQ ID NOS: 7544

; SEQ ID NO 5111

; LENGTH: 53

; TYPE: PRT

; ORGANISM: S.epidermidis

US-09-450-969-5111

Query Match 77.5%; Score 31; DB 3; Length 53;

Best Local Similarity 66.7%; Pred. No. 45;

Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 ILILSKIYV 9

Db 15 ILILTNVYV 23
 ||||: |||

RESULT 6

US-10-724-972B-5111
 ; Sequence 5111, Application US/10724972B
 ; Patent No. 7183083
 ; GENERAL INFORMATION:
 ; APPLICANT: DOUCETTE-STAMM, LYNN
 ; APPLICANT: BUSH, DAVID
 ; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
 ; TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
 ; FILE REFERENCE: 47040.0007US
 ; CURRENT APPLICATION NUMBER: US/10/724,972B
 ; CURRENT FILING DATE: 2003-12-01
 ; PRIOR APPLICATION NUMBER: 09/134,001
 ; PRIOR FILING DATE: 1998-08-13
 ; PRIOR APPLICATION NUMBER: 60/055,779
 ; PRIOR FILING DATE: 1997-08-14
 ; PRIOR APPLICATION NUMBER: 60/064,964
 ; PRIOR FILING DATE: 1997-11-08
 ; NUMBER OF SEQ ID NOS: 7546
 ; SOFTWARE: PatentIn version 3.3
 ; SEQ ID NO 5111
 ; LENGTH: 53
 ; TYPE: PRT
 ; ORGANISM: Staphylococcus epidermidis
 US-10-724-972B-5111

Query Match 77.5%; Score 31; DB 3; Length 53;
 Best Local Similarity 66.7%; Pred. No. 45;
 Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ILILSKIYV 9
 ||||: |||
 Db 15 ILILTNVYV 23

RESULT 7

US-09-270-767-35863
 ; Sequence 35863, Application US/09270767
 ; Patent No. 6703491
 ; GENERAL INFORMATION:
 ; APPLICANT: Homburger et al.
 ; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
 ; FILE REFERENCE: File Reference: 7326-094
 ; CURRENT APPLICATION NUMBER: US/09/270,767
 ; CURRENT FILING DATE: 1999-03-17
 ; NUMBER OF SEQ ID NOS: 62517
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 35863
 ; LENGTH: 167
 ; TYPE: PRT
 ; ORGANISM: Drosophila melanogaster
 US-09-270-767-35863

Query Match 77.5%; Score 31; DB 2; Length 167;
 Best Local Similarity 55.6%; Pred. No. 1.5e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ILILSKIYV 9
 |:::|
 Db 81 IILITKIYV 89

RESULT 8

US-09-270-767-51080
 ; Sequence 51080, Application US/09270767
 ; Patent No. 6703491
 ; GENERAL INFORMATION:
 ; APPLICANT: Homburger et al.
 ; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
 ; FILE REFERENCE: File Reference: 7326-094
 ; CURRENT APPLICATION NUMBER: US/09/270,767
 ; CURRENT FILING DATE: 1999-03-17
 ; NUMBER OF SEQ ID NOS: 62517
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 51080
 ; LENGTH: 167
 ; TYPE: PRT
 ; ORGANISM: Drosophila melanogaster
 US-09-270-767-51080

Query Match 77.5%; Score 31; DB 2; Length 167;
 Best Local Similarity 55.6%; Pred. No. 1.5e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ILILSKIYV 9
 |:::|
 Db 81 IILITKIYV 89

RESULT 9

US-09-248-796A-14687
 ; Sequence 14687, Application US/09248796A
 ; Patent No. 6747137
 ; GENERAL INFORMATION:
 ; APPLICANT: Keith Weinstock et al
 ; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
 ; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
 ; FILE REFERENCE: 107196.132
 ; CURRENT APPLICATION NUMBER: US/09/248,796A
 ; CURRENT FILING DATE: 1999-02-12
 ; PRIOR APPLICATION NUMBER: US 60/074,725
 ; PRIOR FILING DATE: 1998-02-13
 ; PRIOR APPLICATION NUMBER: US 60/096,409
 ; PRIOR FILING DATE: 1998-08-13
 ; NUMBER OF SEQ ID NOS: 28208
 ; SEQ ID NO 14687
 ; LENGTH: 184

; TYPE: PRT
 ; ORGANISM: *Candida albicans*
 US-09-248-796A-14687

Query Match 77.5%; Score 31; DB 2; Length 184;
 Best Local Similarity 55.6%; Pred. No. 1.7e+02;
 Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 ILILSKIYV 9
 : :||||:
 Db 100 LAVLSKIYI 108

RESULT 10

US-10-369-493-7235

; Sequence 7235, Application US/10369493

; Patent No. 7314974

; GENERAL INFORMATION:

; APPLICANT: Cao, Yongwei

; APPLICANT: Hinkle, Gregory J.

; APPLICANT: Slater, Steven C.

; APPLICANT: Goldman, Barry S.

; APPLICANT: Chen, Xianfeng

; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF

; TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES

; FILE REFERENCE: 38-10(52052)B

; CURRENT APPLICATION NUMBER: US/10/369,493

; CURRENT FILING DATE: 2003-02-28

; PRIOR APPLICATION NUMBER: US 60/360,039

; PRIOR FILING DATE: 2002-02-21

; NUMBER OF SEQ ID NOS: 47374

; SEQ ID NO 7235

; LENGTH: 289

; TYPE: PRT

; ORGANISM: *Burkholderia cepacia*

US-10-369-493-7235

Query Match 77.5%; Score 31; DB 3; Length 289;
 Best Local Similarity 55.6%; Pred. No. 2.7e+02;
 Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 ILILSKIYV 9
 :|: ||:|
 Db 213 LLLFSKVYV 221

RESULT 11

US-09-270-767-35094

; Sequence 35094, Application US/09270767

; Patent No. 6703491

; GENERAL INFORMATION:

; APPLICANT: Homburger et al.

; TITLE OF INVENTION: Nucleic acids and proteins of *Drosophila melanogaster*

; FILE REFERENCE: File Reference: 7326-094

; CURRENT APPLICATION NUMBER: US/09/270,767

; CURRENT FILING DATE: 1999-03-17


```
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 35094
; LENGTH: 312
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-35094
```

```
Query Match          77.5%; Score 31; DB 2; Length 312;
Best Local Similarity 55.6%; Pred. No. 2.9e+02;
Matches      5; Conservative      3; Mismatches      1; Indels      0; Gaps      0;
```

```
Qy      1 ILILSKIYV 9
        ::|: ||||
Db      287 VIIIKKIYV 295
```

RESULT 12

```
US-09-270-767-50311
; Sequence 50311, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 50311
; LENGTH: 312
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-50311
```

```
Query Match          77.5%; Score 31; DB 2; Length 312;
Best Local Similarity 55.6%; Pred. No. 2.9e+02;
Matches      5; Conservative      3; Mismatches      1; Indels      0; Gaps      0;
```

```
Qy      1 ILILSKIYV 9
        ::|: ||||
Db      287 VIIIKKIYV 295
```

RESULT 13

```
US-10-369-493-4476
; Sequence 4476, Application US/10369493
; Patent No. 7314974
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
```

```

; APPLICANT: Slater, Steven C.
; APPLICANT: Goldman, Barry S.
; APPLICANT: Chen, Xianfeng
; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
; TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
; CURRENT FILING DATE: 2003-02-28
; PRIOR APPLICATION NUMBER: US 60/360,039
; PRIOR FILING DATE: 2002-02-21
; NUMBER OF SEQ ID NOS: 47374
; SEQ ID NO 4476
; LENGTH: 312
; TYPE: PRT
; ORGANISM: Burkholderia fungorum
US-10-369-493-4476

```

```

Query Match          77.5%; Score 31; DB 3; Length 312;
Best Local Similarity 55.6%; Pred. No. 2.9e+02;
Matches      5; Conservative      3; Mismatches      1; Indels      0; Gaps      0;

```

```

Qy      1 ILILSKIYV 9
       :|: ||:|
Db      232 LLLFSKVYV 240

```

RESULT 14

```

US-10-149-310-260
; Sequence 260, Application US/10149310
; Patent No. 7229784
; GENERAL INFORMATION:
; APPLICANT: Holtzman, Douglas
; APPLICANT: Madden, Kevin T.
; APPLICANT: Maxon, Mary
; APPLICANT: Sherman, Amir
; TITLE OF INVENTION: Modulation of Secondary Metabolite Production by
; TITLE OF INVENTION: Zinc Binuclear Cluster Proteins
; FILE REFERENCE: 14184-019US1
; CURRENT APPLICATION NUMBER: US/10/149,310
; CURRENT FILING DATE: 2003-02-19
; PRIOR APPLICATION NUMBER: PCT/US01/29288
; PRIOR FILING DATE: 2001-09-19
; PRIOR APPLICATION NUMBER: US 60/233,564
; PRIOR FILING DATE: 2000-09-19
; NUMBER OF SEQ ID NOS: 308
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 260
; LENGTH: 758
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
US-10-149-310-260

```

```

Query Match          77.5%; Score 31; DB 3; Length 758;
Best Local Similarity 75.0%; Pred. No. 7.3e+02;
Matches      6; Conservative      2; Mismatches      0; Indels      0; Gaps      0;

```

Qy 1 ILILSKIY 8
 |||:|:|
 Db 409 ILIMSRIY 416

RESULT 15

US-09-248-796A-20980

; Sequence 20980, Application US/09248796A

; Patent No. 6747137

; GENERAL INFORMATION:

; APPLICANT: Keith Weinstock et al

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS

; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: 107196.132

; CURRENT APPLICATION NUMBER: US/09/248,796A

; CURRENT FILING DATE: 1999-02-12

; PRIOR APPLICATION NUMBER: US 60/074,725

; PRIOR FILING DATE: 1998-02-13

; PRIOR APPLICATION NUMBER: US 60/096,409

; PRIOR FILING DATE: 1998-08-13

; NUMBER OF SEQ ID NOS: 28208

; SEQ ID NO 20980

; LENGTH: 883

; TYPE: PRT

; ORGANISM: Candida albicans

US-09-248-796A-20980

Query Match 77.5%; Score 31; DB 2; Length 883;

Best Local Similarity 75.0%; Pred. No. 8.6e+02;

Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ILILSKIY 8
 :|||:|
 Db 65 LLILSRIY 72

Search completed: March 17, 2009, 05:04:35

Job time : 1.76252 secs

SCORE: 0